



IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicant(s): **BOYLE, et al.**

Serial No.: **09/613,591**

Group Art Unit No.: **1647**

Filed: **JULY 10, 2000**

Examiner: **R. Deberry**

For: **COMBINATION THERAPY FOR CONDITIONS
LEADING TO BONE LOSS**

Docket No.: **A-378CIP5**

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RESPONSE AND AMENDMENT

Assistant Commissioner for Patents
Washington, DC 20231

Sir:

In response to the Office Action dated June 4, 2002, Paper No. 25, the Applicants request that the following amendments be made to the above application:

SPECIFICATION

At page 5, lines 14-36, replace with the following paragraph:

Figure 1. A. FASTA analysis of novel EST LORF. Shown is the deduced FRI-1 amino acid sequence aligned to the human TNFR-II sequence (SEQ ID NO: 169 and SEQ ID NO: 138). B. Profile analysis of the novel EST LORF shown is the deduced FRI-1 amino acid sequence aligned to the TNFR-profile (SEQ ID NO: 170 and SEQ ID NO: 178). C. Structural view of TNFR superfamily indicating region which is homologous to the novel FRI-1.

Figure 2. Structure and sequence of full length rat OPG gene, a novel member of the TNFR superfamily. A. Map of pMOB-B1.1 insert. Box indicates position of LORF within the cDNA sequence (bold line). Black box indicates signal peptide, and gray ellipses indicate position of cysteine-rich repeat sequences. B, C. Nucleic acid and protein sequence of the Rat OPG cDNA. The predicted signal peptide is underlined, and potential sites of N-linked glycosylation are indicated in bold, underlined letters (SEQ ID NO: 120 and 121). D, E. Pileup sequence comparison (Wisconsin GCG Package, Version 8.1) of OPG with other members of the TNFR superfamily, fas (SEQ ID NO:128); tnfr1 (SEQ ID NO: 129); sfu-t2 (SEQ ID NO:130); tnfr2 (SEQ ID NO:131); cd40 (SEQ ID NO:132); osteo (SEQ ID NO:133); ngfr (SEQ ID NO:134); ox40 (SEQ ID NO:135); 41bb (SEQ ID NO:136)..

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